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Sheet	1	of	3	Application Number	Not yet assigned
				Filing Date	Herewith
				First Named Inventor	Steven D. Schwartz
				Art Unit	Not yet assigned
				Examiner Name	Not yet assigned
				Attorney Docket Number	96700/855

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	2	BAGDASSARIAN et al., Molecular Electrostatic Potential Analysis for Enzymatic Substrates, Competitive Inhibitors, and Transition-State Inhibitors. <i>J. Am. Chem. Soc.</i> , 118:8825-36, 1996.	
	3	BETTS et al., Cytidine Deaminase. The 2-3 Angstrom Crystal Structure of an Enzyme: Transition-state Analog Complex. <i>J. Mol. Biol.</i> , 235:635-56, 1994.	
	4	BOHM, New Approaches in Molecular Structure Prediction. <i>Biophysical Chemistry</i> , 59:1-32, 1996.	
	5	BRUSIC et al., Prediction of MHC Class II-Binding Peptides Using an Evolutionary Algorithm and Artificial Neural Network. <i>Bioinformatics</i> , 14:121-30, 1998.	
	6	EHRLICH and SCHRAMM, Electrostatic Potential Surface Analysis of the Transition State for AMP Nucleosidase and for Formycin 5'-Phosphate, a Transition-State Inhibitor. <i>Biochem.</i> , 33:8890-96, 1994.	
	7	FRICK et al., Binding of Pyrimidin-2-one Ribonucleoside by Cytidine Deaminase as the Transition-State Analogue 3,4-Dihydrouridine and the Contribution of the 4-Hydroxyl Group to Its Binding Affinity. <i>Biochemistry</i> , 28:9423-30, 1989.	
	8	GASTEIGER et al., Representation of Molecular Electrostatic Potentials by Topological Feature Maps. <i>J. Am. Chem. Soc.</i> , 116:4608-20, 1994.	
	9	HORENSTEIN and SCHRAMM, Electronic Nature of the Transition State for Nucleoside Hydrolase. A Blueprint for Inhibitor Design. <i>Biochemistry</i> , 32:7089-97, 1993.	
	10	KLINE and SCHRAMM, Electrostatic Potential Surfaces of the Transition State for AMP Deaminase and for (R)-Coformycin, a Transition State Inhibitor. <i>J. Biol. Chem.</i> , 269:22385-90, 1994.	
	11	SO and RICHARDS, Application of Neural Networks: Quantitative Structure-Activity Relationships of the Derivatives of 2,4-Diamino-5-(substituted-benzyl) pyrimidines as DHFR Inhibitors. <i>J. Med. Chem.</i> , 35:3201-7, 1992.	

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	12	WAGENER et al., Autocorrelation of Molecular Surface Properties for Modeling Corticosteroid Binding Globulin and Cytosolic Ah Receptor Activity by Neural Networks. J. Am. Chem. Soc., 117:7769-75, 1995.		
	13	WEINSTEIN et al., Predictive Statistics and Artificial Intelligence in the U.S. National Cancer Institute's Drug Discovery Program for Cancer and AIDS. Stem Cells, 12:13-22, 1994.		

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